Title: Genes Encoding Sulfate Assimilation Proteins Inventor: Stephen M. Allen et al. Application No.: To be assigned Docket No.: BB1167US DIV2

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TRKRKYPTRFDTLFPFFSRRKPRAKAPTPTPVMSAAVGGAEFHGFRGGGGGAAQLQRSRM MTTSFGPATTSAPLKDHKVQIPSFHGLRSSSASALPRNA	TISFGAAINIAVADDPNPKLQIHNFSGLKSTSNSLLLSRR	120 LGRPLRVATPHAAAPAGGGGSSSASIRAVSAPLKKDASEVKRSKVEIIKEKSNFLRYPLN LSLPSSTRSLSLIRAVSTPAQSETATVKRSKVEIFKEQSNFIRYPLN	LGRPIRVAPPAAARPGGASAGSIRAVSAPAKKDASEVKRSKVEIIKEKSNFLRYPLN LHVFQSFSPSNPSSIVRAVSTPAKPAAVEPKRSKVEIFKEQSNFIRYPLN	121 EELVSEAPNINDSAVQLIKFHGSYQQTDRDVRGQKNYSFMLRTKNPCGKVPNQLYLAMDT EDILTDAPNISEAATQLIKFHGSYQQYNREERGSRSYSFMIRTKNPCGKVSNQLYLTMDD	EELVSEAPNINESAVQLIKFHGSYQQTDRDVRGQKNYSFMLRTKNPCGKVPNQLYLAMDT EEILNDAPNINEAATQLIKFHGSYMQYDRDERGGRSYSFMLRTKNPGGEVPNRLYLVMDD	240 LADEFGIGTLRLTTRQTFQLHGVLKKNLKTVISTVIKNMGSSLGACGDLNRNVLAPAAPY LADQFGIGTLRLTTRQTFQLHGVLKKDLKTVMGTIIRNMGSTLGACGDLNRNVLAPAAPL	LADEFGIGTLRLTTRQTFQLHGVLKKNLKTVLSTVIKNMGSTLGACGDLNRNVLAPAAPY LADQFGIGTLRLTTRQTFQLHGVLKKNLKTVMSTIIKNMGSTLGACGDLNRNVLAPAAPF
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300 VRKDILFAQETAENIAALLTPQSGAYYDLWVDGEKIMSAEEPPEVTKARNDNTYGTNFPD ARKDYLFAQQTAENIAALLAPQSGFYYDIWVDGEKILTSE-PPEVVQARNDNSHGTNFPD	361 SPEPIYGTQYLPRKFKIAVTVAGDNSVDILTNDIGVVVVSDSAGEPVGFNIYVGGGMGRT SPEPIYGTQFLPRKFKIAVTVPTDNSVDILTNDIGVVVVTDDDGEPQGFNIYVGGGMGRT	420 HRVETTFPRLADPLGYVPKEDILYAIKAIVVTQRENGRRDDRRYSRMKYLIDNWGIEKFR HRLETTFPRLAEPIGYVPKEDILYAVKAIVVTQRENGRRDDRKYSRLKYLISSWGIEKFR	421 AEVEKYYGKKFEDSRPLPEWQFNSYLGWQEQGDGKLFYGVHVDNGRVAGQAKKTLREIIE SVVEQYYGKKFEPFRALPEWEFKSYLGWHEQGDGKLFYGLHVDNGRIGGNMKKTLREIIE AEAEKYYGKKFEDFRPLPEWQFNSYLGWQEQGDGKLFYGVHVDNGRLGGQAKKTLREIIE AEVEKYYGKKFESFRPLPEWQFNSYLGWQEQGDGKLFYGVHVDNGRVGGQAKKTLREIIE SVTEQYYGKKFQPCRELPEWEFKSYLGWHEAGDGSLFCGLHVDNGRVKGAMKKALREIIE
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KYNLEVSITPNONLILCGIDQAWKDPITAALAQSGLLEPKDVDPLNITSMACPALPLCPL KYNLNVRITPNONIILTDVRAAWKRPITTTLAQAGLLQPRFVDPLNITAMACPAFPLCPL		KYNLDVSITPNQNLILCGIDQAWREPITTALAQAGLLEPKDVDPLNLTAMACPALPLCPL	KYNLNVRLTPNQNIILCNIRQAWKRPITTVLAQGGLLQPRYVDPLNLTAMACPAFPLCPL	541	AQTEAERGILPILKRIRAVFDKVGIKDHESVVVRITGCPNGCARPYMAEVGFVGDGPNSY	AITEAERGI PNILKRIRDV FDKVGLKYSESVVVRITGCPNGCARPYMAELGLVGDGPNSY		AQTEAERGILPILKRIRAVFNKVGIKDSESVVVRITGCPNGCARPYMAELGFVGDGPKSY	AITEAERGIPDILKRVRAIFERVGLKYSESVVIRITGCPNGCARPYMAELGLVGDGPNSY	601	QIWLGGTPNQSTLAETFMNKVKLQDIEKVLEPLFSYWNSTRQEGESFGSFTRRTGFDKLK	_	-		QIWLGGTPNQTSLAKTFKDKLKVQDLEKVLEPLFFHWRRKKQSKESFGDFTNRMGFEKLG	
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